RAW SEQUENCE LISTING

re-run

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/520,741
Source: /FWP
Date Processed by STIC: 4/14/06

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IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/520,741   DATE: 10/03/2006
                                             TIME: 08:32:47

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\10032006\J520741.raw

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 3 <110> APPLICANT: Sema ApS
 4          Albrechtsen, Morten
 5          Christensen, Claus
 6          Lukanidin, Eugene
 7          Olesen, Ole
 9 <120> TITLE OF INVENTION: Use of compounds capable of inhibiting the proteolytic
processing of
10          semaphorins for prevention, treatment, diagnosis and prognosis of an invasive
11          disease
13 <130> FILE REFERENCE: P682 PC00
C--> 15 <140> CURRENT APPLICATION NUMBER: US/10/520,741
C--> 15 <141> CURRENT FILING DATE: 2005-01-11
15 <150> PRIOR APPLICATION NUMBER: PCT/DK03/00483
16 <151> PRIOR FILING DATE: 2003-07-11
18 <160> NUMBER OF SEQ ID NOS: 30
20 <170> SOFTWARE: PatentIn version 3.1
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 775
24 <212> TYPE: PRT
25 <213> ORGANISM: Mus sp.
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33 Leu Glu Leu Trp Thr Pro Gly His Ser Ala Asn Pro Ser Tyr Pro Arg
34              20                  25                  30
37 Leu Arg Leu Ser His Lys Glu Leu Leu Glu Leu Asn Arg Thr Ser Ile
38          35                  40                  45
41 Phe Gln Ser Pro Leu Gly Phe Leu Asp Leu His Thr Met Leu Leu Asp
42      50                  55                  60
45 Glu Tyr Gln Glu Arg Leu Phe Val Gly Gly Arg Asp Leu Val Tyr Ser
46  65                  70                  75                  80
49 Leu Asn Leu Glu Arg Val Ser Asp Gly Tyr Arg Glu Ile Tyr Trp Pro
50                  85                  90                  95
53 Ser Thr Ala Val Lys Val Glu Glu Cys Ile Met Lys Gly Lys Asp Ala
54              100                 105                 110
57 Asn Glu Cys Ala Asn Tyr Ile Arg Val Leu His His Tyr Asn Arg Thr
58          115                 120                 125
61 His Leu Leu Thr Cys Ala Thr Gly Ala Phe Asp Pro His Cys Ala Phe
62          130                 135                 140
65 Ile Arg Val Gly His His Ser Glu Glu Pro Leu Phe His Leu Glu Ser
66  145                 150                 155                 160
69 His Arg Ser Glu Arg Gly Arg Gly Arg Cys Pro Phe Asp Pro Asn Ser
70                  165                 170                 175
73 Ser Phe Val Ser Thr Leu Val Gly Asn Glu Leu Phe Ala Gly Leu Tyr
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 74                  180                    185                    190
 77 Ser Asp Tyr Trp Gly Arg Asp Ser Ala Ile Phe Arg Ser Met Gly Lys
 78             195                    200                    205
 81 Leu Gly His Ile Arg Thr Glu His Asp Asp Glu Arg Leu Leu Lys Glu
 82             210                    215                    220
 85 Pro Lys Phe Val Gly Ser Tyr Met Ile Pro Asp Asn Glu Asp Arg Asp
 86 225                     230                    235                    240
 89 Asp Asn Lys Met Tyr Phe Phe Phe Thr Glu Lys Ala Leu Glu Ala Glu
 90             245                    250                    255
 93 Asn Asn Ala His Thr Ile Tyr Thr Arg Val Gly Arg Leu Cys Val Asn
 94             260                    265                    270
 97 Asp Met Gly Gly Gln Arg Ile Leu Val Asn Lys Trp Ser Thr Phe Leu
 98             275                    280                    285
101 Lys Ala Arg Leu Val Cys Ser Val Pro Gly Met Asn Gly Ile Asp Thr
102             290                    295                    300
105 Tyr Phe Asp Glu Leu Glu Asp Val Phe Leu Leu Pro Thr Arg Asp Pro
106 305                     310                    315                    320
109 Lys Asn Pro Val Ile Phe Gly Leu Phe Asn Thr Thr Ser Asn Ile Phe
110             325                    330                    335
113 Arg Gly His Ala Val Cys Val Tyr His Met Ser Ser Ile Arg Glu Ala
114             340                    345                    350
117 Phe Asn Gly Pro Tyr Ala His Lys Glu Gly Pro Glu Tyr His Trp Ser
118             355                    360                    365
121 Leu Tyr Glu Gly Lys Val Pro Tyr Pro Arg Pro Gly Ser Cys Ala Ser
122             370                    375                    380
125 Lys Val Asn Gly Gly Lys Tyr Gly Thr Thr Lys Asp Tyr Pro Asp Asp
126 385                     390                    395                    400
129 Ala Ile Arg Phe Ala Arg Ile Asp Pro Leu Met Tyr Gln Pro Ile Lys
130             405                    410                    415
133 Pro Val His Lys Lys Pro Ile Leu Val Lys Thr Asp Gly Lys Tyr Asn
134             420                    425                    430
137 Leu Arg Gln Leu Ala Val Asp Arg Val Glu Ala Glu Asp Gly Gln Tyr
138             435                    440                    445
141 Asp Val Leu Phe Ile Gly Thr Asp Thr Gly Ile Val Leu Lys Val Ile
142             450                    455                    460
145 Thr Ile Tyr Asn Gln Glu Thr Glu Trp Met Glu Glu Val Ile Leu Glu
146 465                     470                    475                    480
149 Glu Leu Gln Ile Phe Lys Asp Pro Ala Pro Ile Ile Ser Met Glu Ile
150             485                    490                    495
153 Ser Ser Lys Arg Gln Gln Leu Tyr Ile Gly Ser Ala Ser Ala Val Ala
154             500                    505                    510
157 Gln Val Arg Phe His His Cys Asp Met Tyr Gly Ser Ala Cys Ala Asp
158             515                    520                    525
161 Cys Cys Leu Ala Arg Asp Pro Tyr Cys Ala Trp Asp Gly Ile Ser Cys
162             530                    535                    540
165 Ser Arg Tyr Tyr Pro Thr Gly Ala His Ala Lys Arg Arg Phe Arg Arg
166 545                     550                    555                    560
169 Gln Asp Val Arg His Gly Asn Ala Ala Gln Gln Cys Phe Gly Gln Gln
170             565                    570                    575
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PATENT APPLICATION: US/10/520,741   DATE: 10/03/2006
                                          TIME: 08:32:47
Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\10032006\J520741.raw

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174              580                     585              590
177 Ile Glu Ser Asn Ser Thr Leu Leu Glu Cys Thr Pro Arg Ser Leu Gln
178          595                 600                 605
181 Ala Lys Val Ile Trp Phe Tyr Gln Lys Gly Arg Asp Val Arg Lys Glu
182      610                 615                 620
185 Glu Val Lys Thr Asp Asp Arg Val Val Lys Met Asp Leu Gly Leu Leu
186 625              630                 635                     640
189 Phe Leu Arg Val Arg Lys Ser Asp Ala Gly Thr Tyr Phe Cys Gln Thr
190              645                 650                     655
193 Val Glu His Asn Phe Val His Thr Val Arg Lys Ile Thr Leu Glu Val
194          660                 665                 670
197 Val Glu Glu His Lys Val Glu Gly Met Phe His Lys Asp His Glu Glu
198      675                 680                 685
201 Glu Arg His His Lys Met Pro Cys Pro Pro Leu Ser Gly Met Ser Gln
202      690                 695                 700
205 Gly Thr Lys Pro Trp Tyr Lys Glu Phe Leu Gln Leu Ile Gly Tyr Ser
206 705              710                 715                     720
209 Asn Phe Gln Arg Val Glu Glu Tyr Cys Glu Lys Val Trp Cys Thr Asp
210              725                 730                     735
213 Lys Lys Arg Lys Lys Leu Lys Met Ser Pro Ser Lys Trp Lys Tyr Ala
214              740                 745                 750
217 Asn Pro Gln Glu Lys Arg Leu Arg Ser Lys Ala Glu His Phe Arg Leu
218      755                 760                 765
221 Pro Arg His Thr Leu Leu Ser
222      770             775
225 <210> SEQ ID NO: 2
226 <211> LENGTH: 775
227 <212> TYPE: PRT
228 <213> ORGANISM: Mus sp.
230 <400> SEQUENCE: 2
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236 Leu Glu Leu Trp Thr Gly Gly His Thr Ala Asp Thr Thr His Pro Arg
237          20                  25                  30
240 Leu Arg Leu Ser His Lys Glu Leu Leu Asn Leu Asn Arg Thr Ser Ile
241      35                  40                  45
244 Phe His Ser Pro Phe Gly Phe Leu Asp Leu His Thr Met Leu Leu Asp
245      50                  55                  60
248 Glu Tyr Gln Glu Arg Leu Phe Val Gly Gly Arg Asp Leu Val Tyr Ser
249 65              70                  75                      80
252 Leu Ser Leu Glu Arg Ile Ser Asp Gly Tyr Lys Glu Ile His Trp Pro
253              85                  90                  95
256 Ser Thr Ala Leu Lys Met Glu Glu Cys Ile Met Lys Gly Lys Asp Ala
257          100                 105                 110
260 Gly Glu Cys Ala Asn Tyr Val Arg Val Leu His His Tyr Asn Arg Thr
261      115                 120                 125
264 His Leu Leu Thr Cys Gly Thr Gly Ala Phe Asp Pro Val Cys Ala Phe
265      130             135                 140
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268 Ile Arg Val Gly Tyr His Leu Glu Asp Pro Leu Phe His Leu Glu Ser
269 145             150             155             160
272 Pro Arg Ser Glu Arg Gly Arg Gly Arg Cys Pro Phe Asp Pro Ser Ser
273             165             170             175
276 Ser Phe Ile Ser Thr Leu Ile Gly Ser Glu Leu Phe Ala Gly Leu Tyr
277         180             185             190
280 Ser Asp Tyr Trp Ser Arg Asp Ala Ala Ile Phe Arg Ser Met Gly Arg
281         195             200             205
284 Leu Ala His Ile Arg Thr Glu His Asp Asp Glu Arg Leu Leu Lys Glu
285     210             215             220
288 Pro Lys Phe Val Gly Ser Tyr Met Ile Pro Asp Asn Glu Asp Arg Asp
289 225             230             235             240
292 Asp Asn Lys Val Tyr Phe Phe Phe Thr Glu Lys Ala Leu Glu Ala Glu
293             245             250             255
296 Asn Asn Ala His Ala Ile Tyr Thr Arg Val Gly Arg Leu Cys Val Asn
297         260             265             270
300 Asp Val Gly Gly Gln Arg Ile Leu Val Asn Lys Trp Ser Thr Phe Leu
301     275             280             285
304 Lys Ala Arg Leu Val Cys Ser Val Pro Gly Met Asn Gly Ile Asp Thr
305     290             295             300
308 Tyr Phe Asp Glu Leu Glu Asp Val Phe Leu Leu Pro Thr Arg Asp His
309 305             310             315             320
312 Lys Asn Pro Val Ile Phe Gly Leu Phe Asn Thr Thr Ser Asn Ile Phe
313             325             330             335
316 Arg Gly His Ala Ile Cys Val Tyr His Met Ser Ser Ile Arg Ala Ala
317         340             345             350
320 Phe Asn Gly Pro Tyr Ala His Lys Glu Gly Pro Glu Tyr His Trp Ser
321         355             360             365
324 Val Tyr Glu Gly Lys Val Pro Tyr Pro Arg Pro Gly Ser Cys Ala Ser
325     370             375             380
328 Lys Val Asn Gly Gly Arg Tyr Gly Thr Thr Lys Asp Tyr Pro Asp Asp
329 385             390             395             400
332 Ala Ile Arg Phe Ala Arg Ser His Pro Leu Met Tyr Gln Ala Ile Lys
333             405             410             415
336 Pro Ala His Lys Lys Pro Ile Leu Val Lys Thr Asp Gly Lys Tyr Asn
337         420             425             430
340 Leu Lys Gln Ile Ala Val Asp Arg Val Glu Ala Glu Asp Gly Gln Tyr
341     435             440             445
344 Asp Val Leu Phe Ile Gly Thr Asp Asn Gly Ile Val Leu Lys Val Ile
345     450             455             460
348 Thr Ile Tyr Asn Gln Glu Met Glu Ser Met Glu Glu Val Ile Leu Glu
349 465             470             475             480
352 Glu Leu Gln Ile Phe Lys Asp Pro Val Pro Ile Ile Ser Met Glu Ile
353             485             490             495
356 Ser Ser Lys Arg Gln Gln Leu Tyr Ile Gly Ser Ala Ser Ala Val Ala
357         500             505             510
360 Gln Val Arg Phe His His Cys Asp Met Tyr Gly Ser Ala Cys Ala Asp
361     515             520             525
364 Cys Cys Leu Ala Arg Asp Pro Tyr Cys Ala Trp Asp Gly Ile Ser Cys
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365        530                      535                    540
368 Ser Arg Tyr Tyr Pro Thr Gly Thr His Ala Lys Arg Arg Phe Arg Arg
369 545                 550                    555                    560
372 Gln Asp Val Arg His Gly Asn Ala Ala Gln Gln Cys Phe Gly Gln Gln
373                 565                    570                    575
376 Phe Val Gly Asp Ala Leu Asp Lys Thr Glu Glu His Leu Ala Tyr Gly
377             580                    585                    590
380 Ile Glu Asn Asn Ser Thr Leu Leu Glu Cys Thr Pro Arg Ser Leu Gln
381         595                    600                    605
384 Ala Lys Val Ile Trp Phe Val Gln Lys Gly Arg Glu Thr Arg Lys Glu
385     610                    615                    620
388 Glu Val Lys Thr Asp Asp Arg Val Val Lys Met Asp Leu Gly Leu Leu
389 625                 630                    635                    640
392 Phe Leu Arg Leu His Lys Ser Asp Ala Gly Thr Tyr Phe Cys Gln Thr
393             645                    650                    655
396 Val Glu His Ser Phe Val His Thr Val Arg Lys Ile Thr Leu Glu Val
397         660                    665                    670
400 Val Glu Glu Glu Lys Val Glu Asp Met Phe Asn Lys Asp Asp Glu Glu
401     675                    680                    685
404 Asp Arg His His Arg Met Pro Cys Pro Ala Gln Ser Ser Ile Ser Gln
405     690                    695                    700
408 Gly Ala Lys Pro Trp Tyr Lys Glu Phe Leu Gln Leu Ile Gly Tyr Ser
409 705                 710                    715                    720
412 Asn Phe Gln Arg Val Glu Glu Tyr Cys Glu Lys Val Trp Cys Thr Asp
413             725                    730                    735
416 Arg Lys Arg Lys Lys Leu Lys Met Ser Pro Ser Lys Trp Lys Tyr Ala
417             740                    745                    750
420 Asn Pro Gln Glu Lys Lys Leu Arg Ser Lys Pro Glu His Tyr Arg Leu
421     755                    760                    765
424 Pro Arg His Thr Leu Asp Ser
425     770                    775
428 <210> SEQ ID NO: 3
429 <211> LENGTH: 560
430 <212> TYPE: PRT
431 <213> ORGANISM: Mus sp.
433 <400> SEQUENCE: 3
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436 1               5                  10                     15
439 Leu Glu Leu Trp Thr Pro Gly His Ser Ala Asn Pro Ser Tyr Pro Arg
440             20                     25                     30
443 Leu Arg Leu Ser His Lys Glu Leu Leu Glu Leu Asn Arg Thr Ser Ile
444         35                     40                     45
447 Phe Gln Ser Pro Leu Gly Phe Leu Asp Leu His Thr Met Leu Leu Asp
448     50                     55                     60
451 Glu Tyr Gln Glu Arg Leu Phe Val Gly Gly Arg Asp Leu Val Tyr Ser
452 65                 70                     75                     80
455 Leu Asn Leu Glu Arg Val Ser Asp Gly Tyr Arg Glu Ile Tyr Trp Pro
456             85                     90                     95
459 Ser Thr Ala Val Lys Val Glu Glu Cys Ile Met Lys Gly Lys Asp Ala
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VERIFICATION SUMMARY
PATENT APPLICATION: US/10/520,741   DATE: 10/03/2006
                                          TIME: 08:32:48

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\10032006\J520741.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date